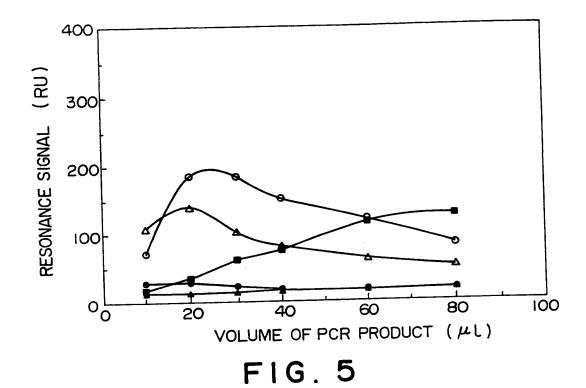


FIG.3

3/6

10 atgaagtgta	20 TATTATTAA		40 TGCCTGTTAC SENSE PRIME		60 TT <u>CGGTATCC</u>
70 TATTCCCGGG SENSE PRIME		90 AGACTTTTCG	100 ACCCAACAAA	110 GTTATGTCTC	120 TTCGTTAAAT
130 AGTATACGGA	140 CAGAGATATC	150 GACCCCTCTT	160 GAACATATA <u>T</u>	170 CTCAGGGGAC SENSE PRI	
	200 ACCACACCCC SENSE PRIM		220 TATTTTGCTG	230 TGGATATACG	240 AGGGCTTGAT
250 GTCTATCAGG	260 CGCGTTTTGA	270 CCATCTTCGT	280 CTGATTATTG	290 AGCAAAATAA	300 TTTATATGTG
	320 TTAATACGGC	330 AACAAATACT	340 TTCTACCGTT		
370	380	390 GGTTTCCATO) 400 G ACAACGGACA	GCAGTTATAC	
) 440 G CGCTGGAACI ENSE PRIME	TTCCGGAAT	D 460 G CAAATCAGTO) 480 GGTTTCATCA
491	n 50	0 51			0 540 C CAGAGCAGTT



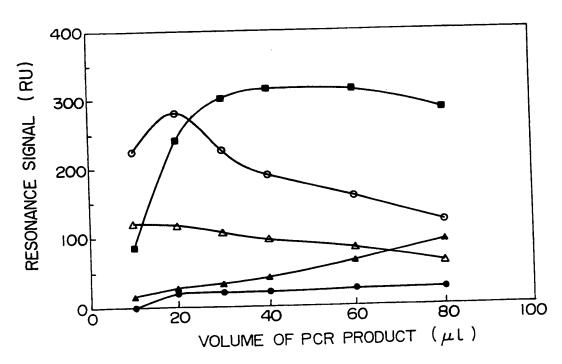
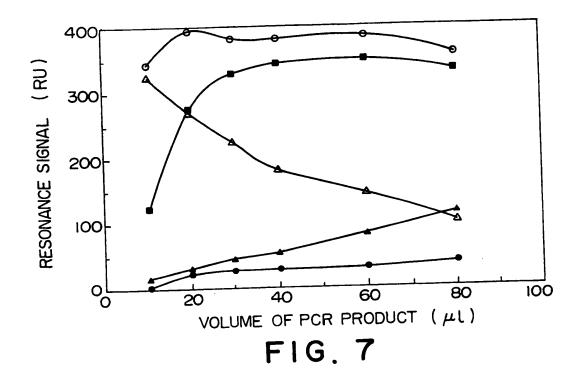
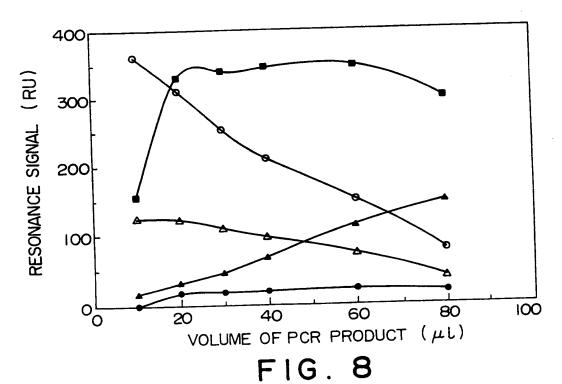


FIG. 6





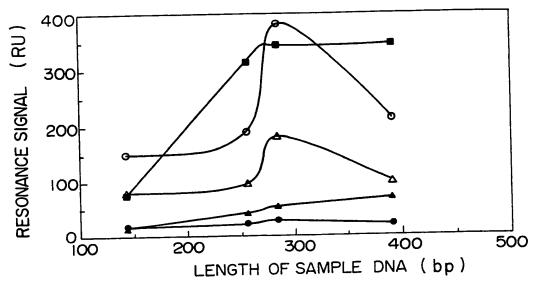


FIG. 9

